OIPE

RAW SEQUENCE LISTING DATE: 12/13/2001 PATENT APPLICATION: US/10/002,854 TIME: 10:27:57

Input Set : A:\M07657038.txt

Output Set: N:\CRF3\12132001\I002854.raw

Does Not Comply
Corrected Diskette Needed

```
4 <110> APPLICANT: Scadden, David T.
              Poznansky, Mark C.
      6
              Olszak, Ivona T.
      7
              Brown, Edward M.
      9 <120> TITLE OF INVENTION: THE CAR RECEPTOR AS A MEDIATOR OF
     10
              MIGRATORY CELL CHEMOTAXIS AND/OR CHEMOKINESIS
     13 <130> FILE REFERENCE: M0765/7038/ERG/KA
C--> 15 <140> CURRENT APPLICATION NUMBER: US/10/002,854
C--> 15 <141> CURRENT FILING DATE: 2001-11-01
     15 <150> PRIOR APPLICATION NUMBER: PCT/US00/15440
     16 <151> PRIOR FILING DATE: 2000-06-02
     18 <160> NUMBER OF SEQ ID NOS: 2
     20 <170> SOFTWARE: FastSEQ for Windows Version 3.0
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ERRORED SEQUENCES

308 <210> SEQ ID NO: 2 309 <211> LENGTH: 1078 310 <212> TYPE: PRT 311 <213> ORGANISM: Homo sapiens 313 <220> FEATURE: 314 <221> NAME/KEY: PEPTIDE 315 <222> LOCATION: (1)...(1078) 316 <223> OTHER INFORMATION: Ca-sensing Receptor 318 <400> SEQUENCE: 2 Met Ala Phe Tyr Ser Cys Cys Trp Val Leu Leu Ala Leu Thr Trp His 321 Thr Ser Ala Tyr Gly Pro Asp Gln Arg Ala Gln Lys Lys Gly Asp Ile 322 25 323 Ile Leu Gly Gly Leu Phe Pro Ile His Phe Gly Val Ala Ala Lys Asp 324 40 325 Gln Asp Leu Lys Ser Arg Pro Glu Ser Val Glu Cys Ile Arg Tyr Asn 326 327 Phe Arg Gly Phe Arg Trp Leu Gln Ala Met Ile Phe Ala Ile Glu Glu 328 70 75 329 Ile Asn Ser Ser Pro Ala Leu Leu Pro Asn Leu Thr Leu Gly Tyr Arg 330 90 331 Ile Phe Asp Thr Cys Asn Thr Val Ser Lys Ala Leu Glu Ala Thr Leu 332 105 333 Ser Phe Val Ala Gln Asn Lys Ile Asp Ser Leu Asn Leu Asp Glu Phe 120 335 Cys Asn Cys Ser Glu His Ile Pro Ser Thr Ile Ala Val Val Gly Ala 336 135 337 Thr Gly Ser Gly Val Ser Thr Ala Val Ala Asn Leu Leu Gly Leu Phe 338 339 Tyr Ile Pro Gln Val Ser Tyr Ala Ser Ser Ser Arg Leu Leu Ser Asn

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										170					175	
340					165	_	_,	_	_	170	71	D	3	3	175	TT 3 0
341	Lys	Asn	Gln		Lys	ser	Phe	Leu		Thr	11e	Pro	ASN	Asp	GIU	HIS
342				180					185	~ 1	_	n1		190	3	m
343	Gln	Ala		Ala	Met	Ala	Asp		He	GLu	Tyr	Pne		Trp	ASII	Trp
344			195			_		200		_		_	205	a 1		a 1
345	Val		Thr	Ile	Ala	Ala		Asp	Asp	Tyr	GTA		Pro	Gly	тте	Glu
346		210					215	_			_	220		_		~
347	Lys	Phe	Arg	Glu	Glu		Glu	Glu	Arg	Asp		Cys	He	Asp	Phe	
348	225					230					235					240
349	Glu	Leu	Ile	Ser	Gln	Tyr	Ser	Asp	Glu	Glu	Glu	Ile	Gln	His		Val
350					245					250					255	
351	Glu	Val	Ile	Gln	Asn	Ser	Thr	Ala	Lys	Val	Ile	Val	Val	Phe	Ser	Ser
352				260					265					270		
353	Gly	Pro	Asp	Leu	Glu	Pro	Leu	Ile	Lys	Glu	Ile	Val	Arg	Arg	Asn	Ile
354	_		275					280					285			
355	Thr	Gly	Lys	Ile	Trp	Leu	Ala	Ser	Glu	Ala	Trp	Ala	Ser	Ser	Ser	Leu
356		290	-		_		295					300				
357	Ile	Ala	Met	Pro	Gln	Tyr	Phe	His	Val	Val	Gly	Gly	Thr	Ile	Gly	Phe
358	305					310					315					320
359	Ala	Leu	Lys	Ala	Gly	Gln	Ile	Pro	Gly	Phe	Arg	Glu	Phe	Leu	Lys	Lys
360					325				-	330	-				335	
361	Va1	His	Pro	Ara	Lvs	Ser	Val	His	Asn	Gly	Phe	Ala	Lys	Glu	Phe	\mathtt{Trp}
362				340		,			345	•			-	350		
363	Glu	Glu	Thr		Asn	Cvs	His	Leu	Gln	Glu	Gly	Ala	Lys	Gly	Pro	Leu
364	014	OLU	355			010		360			- 1		365	•		
365	Dro	Va l		Thr	Phe	Leu	Ara		His	Glu	Glu	Ser	Glv	Asp	Arq	Phe
366	110	370	11DP	1111	1110	200	375	1				380	- 1	•	~	
367	Cor		Ser	Cer	Thr	Δla		Ara	Pro	Leu	Cvs		Glv	Asp	Glu	Asn
368	385	non	OCI	UCL	7111	390	1110	*** 9			395		1			400
369		Cor	Car	Wal	Glu		Pro	Tur	Tle	Asp		Thr	His	Leu	Ara	
370	116	261	361	val	405	1111	110	1 1 1	110	410	-1-	1111		204	415	
371	Cor	TF T T	λαn	บรา		T All	Δla	Va 1	ጥኒኒዮ		T1e	Δla	His	Ala		Gln
372	261	тут	ASII	420	ıyı	цси	niu	· u ·	425	501				430		
373	7 an	Tlo	Пттъ		Cvc	LOU	Dro	G1v		Glv	T.Au	Phe	Thr	Asn	Glv	Ser
	ASP	116	435	TIIT	Cys	цеu	FIO	440	Arg	Ory	ьси	1110	445	11011	0+1	001
374	0	7 1 a		т1.	T *** C	T 770	Wa 1		λΊз	Trn	Gln	Va 1		Lys	His	T.eu
375	Cys		ASP	TTG	ьуѕ	гÃ2	455	GIU	Ala	115	GIII	460	пси	цуз	1115	шеа
376	_	450	.	3	nh -	m la sa		7	Wo+	C1**	C1.,		17 - 1	Пhr	Dho	λcn
377		HIS	ьeu	Asn	Pne		ASII	ASII	Met	GIA	475	GIII	val	Thr	FIIE	480
378	465	_	-1	_	.	470	a1	3	m	C = m		т1о	7 on	m~~	Hic	-
379	GIu	Cys	GTÀ	Asp		val	GIÀ	ASII						Trp		
380	_	_		_	485	_	~ 1 .			490						
381	Ser	Pro	Glu		Gly	ser	TT6	vaı		ьys	GLU	AgT	GTÄ		туг	Asn
382				500				_	505	_,		_	a 1	510		-1 -
383	Val	Tyr		Lys	Lys	GТĀ	GLu		Leu	иne	тте	ASN		Glu	ьys	тте
384			515	_		_	_	520		_	٠,	~	525	a .	0 -	7
385	Leu	_	Ser	Gly	Phe	Ser		GLu	Val	Pro	Phe		Asn	Cys	ser	Arg
386		530					535	_				540	~ 3	~ -	-	m1
387		Cys	Leu	Ala	Gly		Arg	Lys	Gly	Ile		GLu	GLY	GLu	Pro	Thr
388	545					550					555					560

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/002,854

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Input Set : A:\M07657038.txt

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389 390	Cys	Cys	Phe	Glu	Cys 565		Glu	Cys	Pro	Asp 570		Glu	Tyr	Ser	Asp 575	Glu
391	Thr	Asp	Ala				Asn	Lys		Pro	Asp	Asp	Phe	_	Ser	
392	<i>α</i> 1	7	TT	580	C	Q	T1.	77-	585		T1.	a 1	D1-	590		(T)
393 394	GIU	ASI	595		ser	Cys	тте	600		GIU	Ile	GIU	605		Ser	Trp
395	Thr	Glu			Glv	Tle	Δla			T.e.ii	Phe	Δla			C1v	Tlo
396		610		1 110	O ₁	110	615		1111	БСи	1110	620		пси	GLY	116
397	Phe	Leu	Thr	Ala	Phe	Val			Val	Phe	Ile			Ara	Asn	Thr
398	625					630		_			635	_				640
399	Pro	Ile	Val	Lys	Ala	Thr	Asn	Arg	Glu	Leu	Ser	Tyr	Leu	Leu	Leu	Phe
400					645					650		,			655	
401	Ser	Leu	Leu		Cys	Phe	Ser	Ser			Phe	Phe	Ile	Gly	Glu	Pro
402			_	660					665			_		670		
403	Gln	Asp		Thr	Cys	Arg	Leu		Gln	Pro	Ala	Phe		Ile	Ser	Phe
404	17 n 1	т о	675	T] _		O	т1.	680	17- 7	T	m1	3	685	** - 1	.	-
405 406	val	690	Cys	тте	ser	Cys	695	ьeu	vaı	ьys	Thr	700	Arg	vaı	ьeu	Leu
407	Val		Glu	Δla	T.vg	Tle		Thr	Ser	Dhe	His		Lve	Ψrn	Trn	Glv
408	705	1110	014		275	710	110	1111	DCI	1 110	715	nry	цуз	115	115	720
409		Asn	Leu	Gln	Phe		Leu	Val	Phe	Leu	Cys	Thr	Phe	Met	Gln	
410					725					730	•				735	
411	Val	Ile	Cys	Val	Ile	Trp	Leu	Tyr	Thr	Ala	Pro	Pro	Ser	Ser	Tyr	Arg
412				740					745					750		
413	Asn	Gln		Leu	Glu	Asp	Glu		Ile	Phe	Ile	Thr	_	His	Glu	Gly
414	a		755		-	~1	_,	760	7		_		765	_	_	
415	ser		мет	Ala,	Leu	GLY		Leu	TTE	GLY	Tyr		Cys	Leu	Leu	Ala
416 417	λla	770	Cvc	Dho	Dho	Dho	775	Dho	Lva	Cor	Arg	780	Ton	Dro	C1	7 an
418	785	TIE	Cys	FIIE	FILE	790	нта	Pne	пур	ser	795	гу	ьеи	PIO	GIU	800
419		Asn	Glu	Ala	Lvs		Tle	Thr	Phe	Ser	Met	Len	Tle	Phe	Phe	
420					805					810					815	110
421	Val	Trp	Ile	Ser	Phe	Ile	Pro	Ala	Tyr	Ala	Ser	Thr	Tyr	Gly	Lys	Phe
422				820					825					830	_	
423	Val	Ser		Val	Glu	Val	Ile	Ala	Ile	Leu	Ala	Ala	Ser	Phe	Gly	Leu
424	_		835		_ •	_		840			_	_	845			
425	Leu		Cys	Ile	Phe	Phe		Lys	Thr	Tyr	Ile		Leu	Phe	Lys	Pro
426 427	Com	850	3 an	mh so	т1.	a 1	855		7	C	a	860	31-	3	***	. 1 -
427											Ser 875					
429											Arg					
430		 , 5	· uı	mu	885	nrg	AIU	1111	пси	890	пта	SCI	ASII	Val	895	Arg
431	Lys	Arg	Ser	Ser		Leu	Gly	Gly	Ser		Gly	Ser	Thr	Pro		Ser
432	-	_		900			-	_	905					910		
433	Ser	Ile	Ser	Ser	Lys	Ser	Asn	Ser	Glu	Asp	Pro	Phe	Pro	Gln	Pro	Glu
434			915					920					925			
435	Arg		Lys	Gln	Gln	Gln		Leu	Ala	Leu	Thr		Gln	Glu	Gln	Gln
436	a 7	930	_	_	-1	_	935		1		_	940				
437	GIn	Gln	Pro	Leu	Thr	Leu	Pro	Gln	Gln	Gln	Arg	Ser	Gln	Gln	Gln	Pro

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438	945	950	955	960
439	Arg Cys Lys Gln Lys	s Val Ile Phe Gly	Ser Gly Thr Val Thr P	he Ser
440	965	5	970 97	75
441	Leu Ser Phe Asp Glu	u Pro Gln Lys Asn A	Ala Met Ala His Arg As	sn Ser
442	980	985	990	
443	Thr His Gln Asn Se	r Leu Glu Ala Gln 1	Lys Ser Ser Asp Thr Le	eu Thr
444	995	1000	1005	
445	Arg His Gln Pro Leu	u Leu Pro Leu Gln (Cys Gly Glu Thr Asp Le	eu Asp
446	1010	1015	1020	-
447	Leu Thr Val Gln Glu	u Thr Gly Leu Gln (Gly Pro Val Gly Gly As	sp Gln
448	1025	1030	1035	(104) 1040
449	Arg Pro Glu Val Glu	u Asp Pro Glu Glu I	Leu Ser Pro Ala Leu Va	al Val
450	104	45 1	1050 10)55
451	Ser Ser Ser Gln Ser	r Phe Val Ile Ser (Sly Gly Gly Ser Thr Va	al Thr
452	1060	1065	1070	
453	Glu Asn Val Val Asn	n Ser		
454	1075			
	439 440 441 442 443 444 445 446 447 448 449 450 451 452	439 Arg Cys Lys Gln Ly 440 96 441 Leu Ser Phe Asp Gl 442 980 443 Thr His Gln Asn Se 444 995 445 Arg His Gln Pro Le 446 1010 447 Leu Thr Val Gln Gl 448 1025 449 Arg Pro Glu Val Gl 450 1060 451 Ser Ser Ser Gln Se 452 1060 453 Glu Asn Val Val Asi	439 Arg Cys Lys Gln Lys Val Ile Phe Gly 3 440 965 441 Leu Ser Phe Asp Glu Pro Gln Lys Asn 3 442 980 985 443 Thr His Gln Asn Ser Leu Glu Ala Gln 1 444 995 1000 445 Arg His Gln Pro Leu Leu Pro Leu Gln Glu 1 446 1010 1015 447 Leu Thr Val Gln Glu Thr Gly Leu Gln Glu I 448 1025 1030 449 Arg Pro Glu Val Glu Asp Pro Glu Glu I 450 1045 451 Ser Ser Ser Gln Ser Phe Val Ile Ser Glu Glu Asp Glu Asn Val Val Asn Ser	439 Arg Cys Lys Gln Lys Val Ile Phe Gly Ser Gly Thr Val Thr P. 440 965 970 9 441 Leu Ser Phe Asp Glu Pro Gln Lys Asn Ala Met Ala His Arg A. 442 980 985 990 443 Thr His Gln Asn Ser Leu Glu Ala Gln Lys Ser Ser Asp Thr Leu Pro Leu Gln Cys Gly Glu Thr Asp Leu Glo 1010 1015 1020 445 Arg His Gln Pro Leu Leu Pro Leu Gln Cys Gly Glu Thr Asp Leu Glo 1020 447 Leu Thr Val Gln Glu Thr Gly Leu Gln Gly Pro Val Gly Gly Ast 1025 1030 1035 448 1025 1030 1035 449 Arg Pro Glu Val Glu Asp Pro Glu Glu Leu Ser Pro Ala Leu Valon Ser Ser Ser Gln Ser Phe Val Ile Ser Gly Gly Gly Ser Thr Valon 1060 1065 1070 451 Ser Ser Ser Gln Ser Phe Val Ile Ser Gly Gly Gly Ser Thr Valon 1060 1065 1070 453 Glu Asn Val Val Asn Ser

 VERIFICATION SUMMARY
 DATE: 12/13/2001

 PATENT APPLICATION: US/10/002,854
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L:15 M:270 C: Current Application Number differs, Replaced Current Application No

L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:303 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:1

L:448 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:2